

OIPe

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/615,872DATE: 07/25/2000
TIME: 14:33:46Input Set : A:\10312-21.app
Output Set: N:\CRF3\07252000\I615872.raw

ENTERED

3 <110> APPLICANT: AREPALLY, Gowthami
4 KISIEL, Walter
5 KAMEI, Keiko
6 KAMEI, Shintaro
8 <120> TITLE OF INVENTION: Compositions and Methods Useful for the Diagnosis and
9 Treatment of Heparin Induced
10 Thrombocytopenia/Thrombosis
12 <130> FILE REFERENCE: 10312-2U1
C--> 14 <140> CURRENT APPLICATION NUMBER: US/09/615,872
15 <141> CURRENT FILING DATE: 2000-07-13
17 <150> PRIOR APPLICATION NUMBER: US 60/143,536
18 <151> PRIOR FILING DATE: 1999-07-13
20 <160> NUMBER OF SEQ ID NOS: 10
22 <170> SOFTWARE: PatentIn Ver. 2.1
24 <210> SEQ ID NO: 1
25 <211> LENGTH: 120
26 <212> TYPE: PRT
27 <213> ORGANISM: Mus musculus
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33 Ser Val Lys Leu Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Asn Tyr
34 20 25 30
36 Phe Ile Tyr Trp Val Lys Gln Arg Pro Gly Gln Gly Leu Glu Trp Ile
37 35 40 45
39 Gly Glu Ile Asn Pro Arg Asn Gly Asp Thr Asp Phe Asn Glu Lys Phe
40 50 55 60
42 Glu Ser Arg Ala Thr Leu Thr Val Asp Lys Ser Ser Thr Ala Tyr
43 65 70 75 80
45 Met Gln Leu Ser Ser Leu Thr Ser Glu Asp Ser Ala Ile Tyr Tyr Cys
46 85 90 95
48 Thr Arg Ser Pro Tyr Gly Asn Asn Tyr Gly Phe Thr Tyr Trp Gly Gln
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67 Gly Thr Asn Val Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Asn
68 35 40 45
70 Ala Leu Ile Tyr Ser Ala Ser Tyr Arg Tyr Ser Gly Val Pro Asp Arg

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71      50      55      60
73 Phe Thr Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Thr Asn
74 65      70      75      80
76 Val Gln Ser Glu Asp Leu Ala Asp Tyr Phe Cys Gln Gln Tyr Asn Ser
77      85      90      95
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80      100      105      110
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84 <211> LENGTH: 360
85 <212> TYPE: DNA
86 <213> ORGANISM: Mus musculus
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90 tcctgtaagg cttctggcta caccttcacc aattacttta tatactgggt gaagcagagg 120
91 cctggacaag gccttgagtg gattggagag attaatccta gaaatgggtg tactgacttc 180
92 aatgagaagt tcgagagcag ggccacactg actgtagaca aatcctccag cacggcatac 240
93 atgcaactca gcagcctgac atctgaggac tctgcgatct attactgtac aagatcccc 300
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104 agggtcaccg tcacctgcaa ggccagtcag aatgtgggta ctaatgtagc ctgggtatcaa 120
105 cagaaaccag ggcaatctcc taatgcactg atttactcgg catcctaccg gtacagtggg 180
106 gtccctgata gcttcacagg cagtggatct gggacagatt tcactctcac catcaccaat 240
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136 <211> LENGTH: 32
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141 <223> OTHER INFORMATION: Description of Artificial Sequence:PCR Primer
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150 <213> ORGANISM: Artificial Sequence
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VERIFICATION SUMMARY

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L:14 M:270 C: Current Application Number differs, Replaced Current Application Number